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Citation for published version (APA):

Hemker, H. C., Esnouf, M. P., Hemker, P. W., Swart, A. C. W., & MacFarlane, R. G. (1967). Formation of prothrombin converting activity. *Nature*, 215(5098), 248-251. <https://doi.org/10.1038/215248a0>

Document status and date:

Published: 01/01/1967

DOI:

[10.1038/215248a0](https://doi.org/10.1038/215248a0)

Document Version:

Version created as part of publication process; publisher's layout

Please check the document version of this publication:

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Formation of Prothrombin Converting Activity

After measurement of the conversion of prothrombin to thrombin in a purified system, a model has been constructed for the kinetics of the formation of prothrombin converting activity. In the model phospholipid provides a surface on to which coagulant factors bind.

by

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In blood coagulation prothrombin is converted to thrombin as the result of the interaction of several blood coagulation factors^{1,2}. One of the features of blood coagulation theory, which is still controversial, is the precise identity of the prothrombin converting activity. While most authors agree that conversion of prothrombin requires the presence of activated factor X (factor X_a), factor V, phospholipid and calcium ions, the form of their interaction is still in debate. Some authors suggest that the final prothrombin converting substance is a form of factor V modified by its interaction with factor X_a (ref. 3), while others hold the view that factor V facilitates the enzyme conversion of prothrombin to thrombin by factor X_a (ref. 4). The function of the phospholipid in this system seems to depend on its providing a suitable surface on which this reaction may take place, rather than to any specific chemical constitution⁴.

We describe here experiments performed with purified reagents, in which the kinetics of prothrombin activation have been investigated with a view to elucidating further the nature of the process.

Bovine prothrombin was prepared and given by Dr K. Denson⁵. The preparation contained 10,000 NIH μ /ml. of prothrombin, and less than 10 per cent of contaminating material that had neither coagulant nor inhibitory activity. Bovine factor V was prepared according to Esnouf and Jobin⁶; the stock solution contained 40,000 μ /ml. and had

no detectable contamination. Factor X_a was prepared by the conversion of factor X (ref. 7) by the coagulant protein from Russell's viper venom⁸, and then separating the venom protein from factor X_a by chromatography⁴. Factor X_a was not contaminated with any other clotting factors.

The phospholipid was a commercial preparation of mixed phospholipids, 'Inosithin'. Emulsions of this phospholipid were prepared as described by Jobin and Esnouf⁴.

The experiments were carried out as follows. Phospholipid, factor X_a and factor V were added together to the final concentration desired in a medium containing 12 mmolar calcium chloride and 40 mmolar *tris*-hydrochloric acid buffer, pH 7.5. The mixture thus obtained was called the "first incubation mixture". It was kept at 37° C during the "first incubation time". Except in the experiments shown in Fig. 1 where it is plotted along the abscissa, this time was always 4-6 min. After this the first incubation mixture was put in an ice bath at 0° C which prevented appreciable decay of prothrombin converting activity for at least 30 min.

To 1.8 ml. of this "first incubation mixture" 0.2 ml. of the prothrombin solution was added at zero time of the second incubation. The concentration of prothrombin in the second incubation was thus 1,000 NIH u/ml.; this concentration represents an excess of substrate for the prothrombin activator, and did not seem to be inhibitory.

From the second incubation mixture 0.1 ml. portions were sub-sampled into 0.1 ml. of fibrinogen solution of 100 mg/ml. (human fibrinogen Kabi). The clotting time of this mixture was measured in a water bath at 37° C. The clotting times thus obtained were converted into concentrations of thrombin by comparison with the clotting times obtained by dilutions of a standard thrombin solution, in comparable conditions of pH, ionic strength, calcium ion concentration and temperature.

The amount of thrombin thus found was plotted against the incubation time in the second incubation mixture, and from this graph the initial velocity of thrombin formation induced by the prothrombin activator was estimated.

A representative experiment is shown in Fig. 1. In this experiment a mixture with high prothrombin converting activity was diluted 1 in 3 and 2 in 3, and thrombin generation has been followed in the original sample as well as in the dilutions. The two lower lines in the graph are drawn at 2/3 and 1/3 of the slope of the steepest line. It can be seen that: (a) the thrombin generation curve cannot be distinguished from a straight line in the initial phase; (b) the initial velocity of thrombin generation can be regarded as proportional to the amount of prothrombin

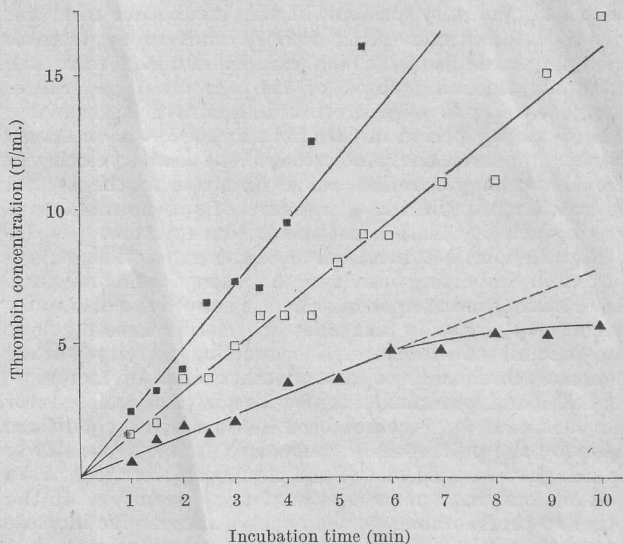


Fig. 1. Thrombin generation in a mixture containing prothrombin and prothrombinase activity. (■—■) Thrombin generation in a mixture containing 20 U/ml. of factor V, 8 U/ml. of factor X_a, 4 γ /ml. of phospholipid, 1,000 NIH U/ml. of prothrombin, 12 mmolar of calcium chloride, and 40 mmolar *tris* hydrochloric acid buffer, pH 7.5. (□—□) Thrombin generation in the same mixture immediately after a 2 in 3 dilution with 40 mmolar *tris*-hydrochloric acid buffer, pH 7.5 containing 12 mmolar calcium chloride. (▲—▲) The same immediately after a 1 in 3 dilution.

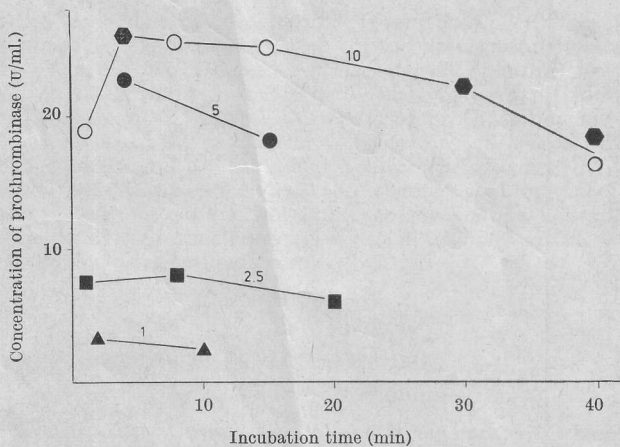


Fig. 2. Generation of prothrombinase activity in a mixture containing factor V, factor X_a and phospholipid. The final concentrations were 25 U/ml. of factor V, 10, 5, 2.5 or 1 U/ml. of factor X_a (as indicated), 5 γ /ml. of phospholipid, 12 mmolar calcium chloride, and 40 mmolar *tris*-hydrochloric acid buffer, pH 7.5. The concentration of prothrombinase is expressed in arbitrary units, one unit being defined as the amount that causes a velocity of thrombin generation of 1 NIH U/10 min during the second incubation.

converting activity present in the incubation mixture; (c) prothrombin converting activity tends to be unstable with larger dilutions, which can be deduced from the gradually declining slope of the thrombin generation curve at a low level of prothrombin converting activity; and (d) no lag period in thrombin formation is observed. It thus appeared acceptable to use the initial velocity of the thrombin generation curve obtained in the second incubation mixture, as a measure of prothrombin converting activity in the first incubation mixture.

Fig. 2 shows a typical curve of the generation of prothrombin converting activity in a first incubation mixture. After mixing the components there is a rapid rise of activity which very gradually falls off again. As will be seen below, the amount of prothrombin converting activity formed bears a relationship to the concentrations of factors V and X_a , and phospholipid. By varying the amounts of the reactants we have not been able to create conditions whereby the prothrombin converting activity rose slowly to a high value. If one of the clotting factors (X_a or V) was an enzyme which catalysed the conversion of the other to prothrombinase, then such a progressive increase in activity in certain conditions might be expected.

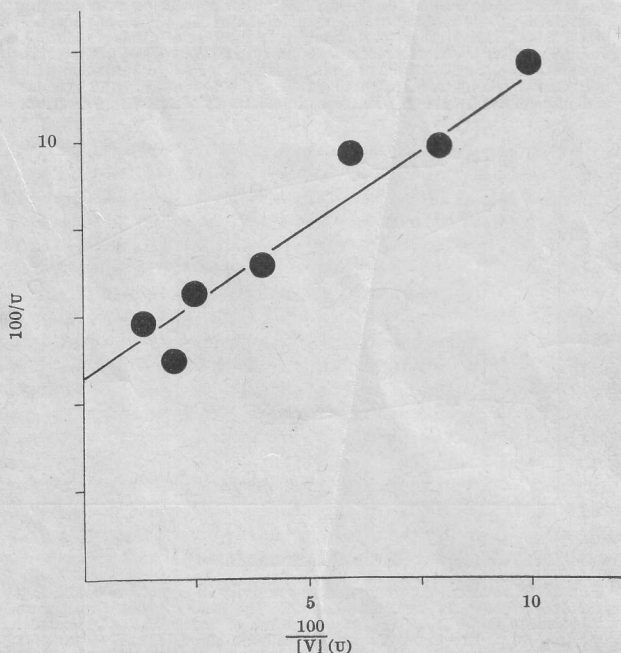
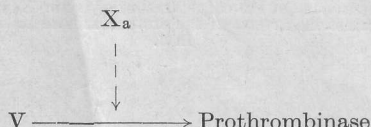


Fig. 3. Relation between concentration of factor V and prothrombinase activity. The inverse of the concentration of factor V expressed in v is plotted against the inverse of the prothrombinase activity expressed in v.

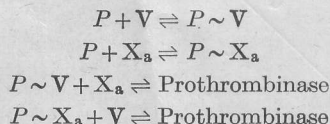
At relatively large concentrations (causing a velocity of thrombin formation of ~ 20 NIH U of thrombin/ml./10 min), the half-life time of the prothrombin converting activity was estimated to be about 55 min; the half-life time of the prothrombin converting activity generated in a thrombin generation test (that is, in a system using normal plasma) was estimated to be about 10 min in the same conditions. Thus the prothrombin converting activity appeared to be more stable in its purified form than it is in plasma. This again emphasizes the point that an inhibitor of prothrombin conversion is likely to be present in plasma⁹.

Figs. 3, 4 and 5 show that a linear relationship between the reciprocal of the concentration of either of the reactants and the reciprocal of the final prothrombin converting activity is the simplest kind of relationship that can be deduced from these experiments. Essentially the same results were obtained when the amount of thrombin was estimated spectrophotometrically as *n*-benzoyl arginine methyl esterase activity.

This relationship cannot be accounted for by the reaction scheme of the type



or by any of its modifications. It can, however, be accounted for by a model based on the assumptions that: (a) there are a fixed number of factor X_a binding sites and a fixed number of factor V binding sites on each unit phospholipid micelle surface; (b) prothrombin conversion occurs when a factor X_a molecule and a factor V molecule, bound to the phospholipid surface, are in a favourable position one to the other; (c) binding of one of the factors does not influence the binding characteristics of the other one⁵. This model can be written in the following reaction formulae



The word prothrombinase refers to the complex which converts prothrombin to thrombin.

With these assumptions the following relationships will exist: (a) the concentration of prothrombinase is proportional to the number of bound factor X_a molecules/unit of phospholipid surface as well as to the amount of bound factor V molecules per unit of phospholipid surface;

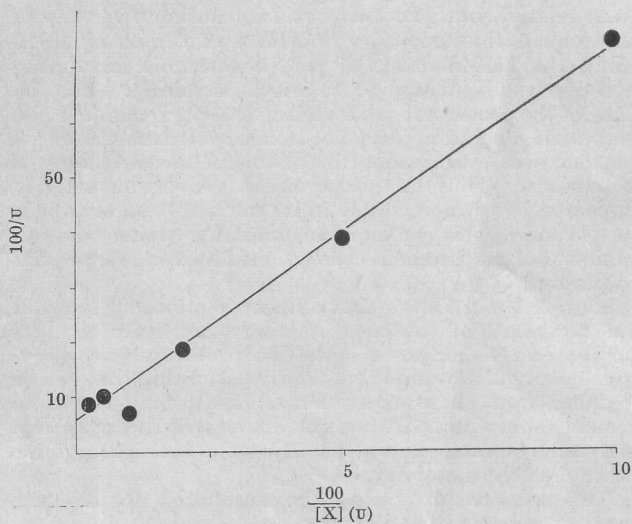


Fig. 4. Relation between concentration of factor X_a and prothrombinase activity. The inverse of the concentration of factor X_a expressed in v is plotted against the inverse of the prothrombinase activity expressed in v .

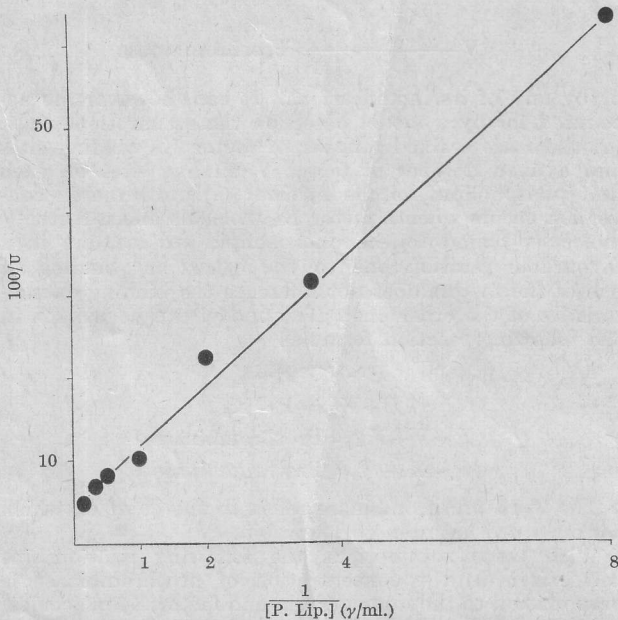


Fig. 5. Relation between phospholipid and prothrombinase activity. The inverse of the phospholipid concentration expressed in $\gamma/ml.$

(b) the concentration of prothrombinase is proportional to the amount of phospholipid surface. This is written in a formula

$$C = K \cdot P \cdot \frac{P \sim X}{P} \cdot \frac{P \sim V}{P} = \frac{K}{P} \cdot P \sim X \cdot P \sim V$$

Where C is prothrombinase concentration; P is phospholipid concentration; $P \sim X$ is phospholipid bound factor X_a concentration; $P \sim V$ is phospholipid bound factor V concentration, and K is a constant. With the aid of elementary mathematics it can be shown

$$\text{from } P + X \xrightleftharpoons[k_{-1}]{k_{+1}} P \sim X; k_{-1}/k_{+1} = K'$$

$$\text{that } \frac{1}{P \sim X} = \frac{1}{X} + \frac{1}{P} + \frac{K'}{X \cdot P} \text{ and}$$

$$\text{from } P + V \xrightleftharpoons[k_{-2}]{k_{+2}} P \sim V; k_{-2}/k_{+2} = K''$$

$$\text{that } \frac{1}{P \sim V} = \frac{1}{V} + \frac{1}{P} + \frac{K''}{V \cdot P}$$

$$\text{So that } \frac{1}{C} = \frac{P}{K} \left(\frac{1}{X} + \frac{1}{P} + \frac{K'}{X \cdot P} \right) \cdot \left(\frac{1}{V} + \frac{1}{P} + \frac{K''}{V \cdot P} \right)$$

In these equations P , X and V indicate the total concentrations of phospholipid, factor X_a and factor V , respectively. For the two protein factors this formula accounts for the relations found: $1/C$ is linearly proportional to $1/V$ at fixed V and P concentrations and linearly proportional to $1/V$ at fixed X and P concentrations.

When the relationship between C and P is examined the formula is best written in the form

$$\frac{1}{C} = P + \frac{1}{P} (X + K') (V + K'') + (V + X + K' + K'') \frac{1}{XVK}$$

At small concentrations of phospholipid P will become negligible compared with $\frac{1}{P} (X + K') (V + K'')$ and the formula will reduce to

$$\frac{1}{C} = \frac{1}{P} (X + K') (V + K'') + (V + X + K' + K'') \frac{1}{XVK}$$

which accounts for the relationship which we found.

At large concentrations of phospholipid the formula will reduce to

$$\frac{1}{C} = P + (V + X + K' + K'') \frac{1}{VXK}$$

which predicts the inhibiting effect of excess phospholipid that is often found.

This inhibitory action of excess phospholipid is not seen in the experiments shown in Fig. 5, because the range of phospholipid tested does not include the pertinent concentrations.

From these results it can be seen that near maximal velocity is often found at relatively small concentrations of the three reactants. This finding explains many of the reports in the literature, stating that further addition of one particular reactant did not increase the final yield of prothrombinase. The conclusion that this reactant may not therefore be a constituent of prothrombinase is apparently not justified.

The present investigations have partly been carried out under the auspices of the Netherlands Foundation for Chemical Research (SON) and with financial support from the Netherlands Organization for the Advancement of Pure Research (ZWO). We thank Mr P. P. Devilee for technical assistance.

Received May 8, 1967.

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